

SCORE Search Results Details for Application 10597173 and Search Result 20070521_150829_us-10-597-173b-7.100align.rge.

Score Home	Retrieve Application	SCORE System	SCORE	Comments /
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This page gives you Search Results detail for the Application 10597173 and Search Result 20070521_150829_us-10-597-173b-7.100align.rge.

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GenCore version 6.2.1
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DM nucleic - nucleic search, using sw model

Run on: May 23, 2007, 00:36:17 ; Search time 817 Seconds
(without alignments)
1268.619 Million cell updates/sec

Title: US-10-597-173B-7
Perfect score: 15
Sequence: 1 ttcttttgaaaactga 15

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 7568541 seqs, 34560148153 residues

Total number of hits satisfying chosen parameters: 15137082

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 300 summaries

Database : GenEmbl:*
1: gb_env:*
2: gb_pat:*
3: gb_ph:*
4: gb_pl:*
5: gb_pr:*
6: gb_ro:*
7: gb_sts:*
8: gb_sy:*

RESULT 44

JRED6PCA

LOCUS URED6PCA 305 bp DNA linear BCT 21-MAY-1999

DEFINITION Ureaplasma canigenitalium (canine ureaplasma) 16S-23S rRNA genes, intergenic spacer region.

ACCESSION D63684

VERSION D63684.1 GI:944983

KEYWORDS 16S rRNA; 23S rRNA.

SOURCE Ureaplasma canigenitalium

ORGANISM Ureaplasma canigenitalium

Bacteria; Firmicutes; Mollicutes; Mycoplasmataceae; Ureaplasma.

REFERENCE 1

AUTHORS Harasawa,R., Lefkowitz,E.J., Glass,J.I. and Cassell,G.H.

TITLE Phylogenetic analysis of the 16S-23S rRNA intergenic spacer regions of the genus Ureaplasma

JOURNAL J. Vet. Med. Sci. 58 (3), 191-195 (1996)

PUBMED 8777224

REFERENCE 2 (bases 1 to 305)

AUTHORS Harasawa,R.

TITLE Direct Submission

JOURNAL Submitted (27-JUL-1995) Ryo Harasawa, The University of Tokyo Faculty of Medicine, Animal Center for Biomedical Research; Hongo 7-3-1, Bunkyo-ku, Tokyo 113-0033, Japan (E-mail:ryo@m.u-tokyo.ac.jp, Tel:+81-3-5841-3623, Fax:+81-3-5841-3679)

FEATURES Location/Qualifiers

source

1. .305

/organism="Ureaplasma canigenitalium"

/mol_type="genomic DNA"

/strain="D6P-C"

/specific_host="canine"

/db_xref="taxon:42092"

misc_feature

1. .305

/note="16S-23S rRNA genes intergenic spacer region"

ORIGIN

Query Match 100.0%; Score 15; DB 15; Length 305;

Best Local Similarity 100.0%; Pred. No. 2.3e+03;

Matches 15; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

2y 1 TTCTTTGAAAACTGA 15

|||||

2b 180 TTCTTTGAAAACTGA 194

RESULT 45

AJ780992

LOCUS AJ780992 305 bp DNA linear BCT 28-JUL-2004

DEFINITION Mycoplasma gallinarum 16S-23S intergenic spacer, type strain PG16T.